

1

2 **Discussion**

3 Tamoxifen performs its anti-tumor activity via many pathways, both ER-dependent
4 and ER-independent. In the present study, we attempted to identify novel anti-tumor
5 compounds among RIDs that were designed from a lead compound, tamoxifen. We
6 synthesized forty-eight RIDs by efficient methods including the three-component
7 coupling reaction⁹). To screen them for promising anti-tumor compounds, we used the
8 JFCR39 panel and an ER binding assay. By this means, we found two RIDs, RID-SB1
9 and RID-SB8, that showed stronger growth-inhibitory activity than tamoxifen and are
10 expected to have unique modes of action.

11 The forty-eight RIDs showed growth-inhibitory activities against JFCR39 over a
12 wide range of MG-MIDs from 0.85 to 43.7 μM . Forty out of forty-eight RIDs (83%)
13 showed higher growth-inhibitory activity than tamoxifen (MG-MID = 7.41 μM). The
14 SAR study indicated that the structures of aminoalkoxyphenyl groups at the C-1
15 position and the common central ethylenic double bond were important in retaining a
16 high level of growth-inhibitory activity.

17 The ER α binding activity of the RIDs was spread across a huge range of IC₅₀ values
18 from 26.7 to > 10000 nM. In general, the 1st-generation RIDs exhibited ER α binding
19 activities. Among them RID-G showed the most potent activity (IC₅₀ = 26.6 nM), which
20 is close to the activity of tamoxifen (IC₅₀ = 21.1 nM). In contrast, a number of RIDs in
21 the 2nd-generation were inactive with respect to ER α binding. The SAR study clearly
22 indicated that both the phenyl and ethyl groups at the C-2 position of the central
23 ethylenic bond were essential for ER α binding activity. In addition, there was a
24 tendency for RIDs bearing more bulky structures around the double bond (RID-SB7 to
25 SB12 and SG7 to SG12) to display higher ER α binding activity.

1 According to the above observations, the forty-eight RID compounds were profiled
2 by growth-inhibitory activity and ER α binding activity (Fig. 5). Interestingly, these two
3 activities didn't correlate in the forty-eight RIDs. Some of them, such as RID-G, showed
4 high levels in both activities, while others were active in only one. In the latter group,
5 we focused on RID-SB1 and RID-SB8 because they were highly active in
6 growth-inhibition but were inactive in regard to ER α binding. It was suggested that they
7 inhibited the cell growth via an ER-independent mechanism and thus may inhibit the
8 growth of cancer cells by a different mode of action than tamoxifen. The COMPARE
9 analysis was therefore performed on this study, and it revealed that both RID-SB1 and
10 RID-SB8 showed very weak correlation coefficients ($r < 0.4$) with tamoxifen. These
11 results support the hypothesis above.

12 So far, a number of tamoxifen derivatives have been designed and synthesized with
13 the aim of enhancing ER binding activity and reducing side-effects. However, this has
14 not been entirely successful. For example, clomifene²⁴⁾, ospemifene^{25,26)}, iodoxifene²⁷⁾,
15 raloxifene²⁷⁾, arzoxifene²⁸⁾, lasofoxifene²⁹⁾ and levormeloxifene^{30,31)}, possess high ER
16 binding activity, while still having such disadvantages as biotransformation effect,
17 biological isomerization, thromboembolic effect and agonist action in uterus³²⁾. Most of
18 these tamoxifen derivatives show very similar effects as tamoxifen and have been
19 proposed considering only the effect on ER. In the present study, RID-SB1 and
20 RID-SB8, while having completely lost ER binding activity, display approximately
21 5-fold higher antitumor activity than tamoxifen, indicating a non-ER target in its
22 anti-tumor process. Such non-ER function might be useful in unveiling new targets to
23 inhibit tumor cell proliferation. Therefore, RID-SB1 and RID-SB8 maybe novel drug
24 candidates for breast cancer but also for other malignancies.

1 We finally attempted to predict the action modes of RID-SB1 and RID-SB8 by using
2 our JFCR39 drug database and COMPARE analysis. The two RIDs did not correlate
3 with any currently available anti-cancer drug, suggesting that they could be examples of
4 a novel class of anti-cancer drug. The COMPARE analysis suggested some targets
5 shown in Table 2, however, this needs to be verified by biological testing in the future.

6 In conclusion, we synthesized forty-eight tamoxifen-derivatives, RIDs, and screened
7 them, searching for novel candidates of a new class of anti-cancer drug. We identified
8 RID-SB1 and RID-SB8 as having potent tumor growth-inhibitory activity but having
9 completely lost ER α binding activity. Based on these results and COMPARE analysis, it
10 was suggested that RID-SB1 and RID-SB8 had unique action modes, different from
11 those of current anti-cancer drugs including tamoxifen. RID-SB1 and RID-SB8 merit
12 further investigation.

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11

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- 6

1 **Figure legends**

2 Fig. 1. Chemical Structures of tamoxifen, 4-hydroxytamoxifen and RIDs. (A) tamoxifen,
3 4-hydroxytamoxifen and 1st-generation RIDs. (B) 2nd-generation RIDs. (C)
4 3rd-generation RIDs.

5 Fig. 2. Dose response curves of (A) tamoxifen and (B) RID-SB1 against growth of
6 JFCR39 cancer cell lines. The growth inhibition of tamoxifen and RID-SB1 on 39 cell
7 lines (Br, breast; CNS, central nervous system; Co, colorectal; Lu, lung; Me, melanoma;
8 Ov, ovarian; Re, renal; St, stomach; xPg, prostate) were measured as described in
9 Materials and methods.

10 Fig. 3. Fingerprints of tamoxifen, RID-SB1 and RID-SB8. Fingerprint shows the
11 differential growth inhibition pattern of chemicals against JFCR 39 cancer cell lines.
12 The *X*-axis represents difference in logarithmic scale between mean of LogGI₅₀ values
13 for 39 cell lines and the LogGI₅₀ for each cell line. Bars to the right of 0 indicates cell
14 lines that are sensitive to the compound, in contrast, bars on the left of 0 means the
15 resistance. MG-MID, mean of LogGI₅₀ values for 39 cell lines; Delta, difference
16 between the MG-MID and the Log GI₅₀ value for the most sensitive cell line; Range,
17 difference between the LogGI₅₀ values for the most resistant cell and the most sensitive
18 cell line.

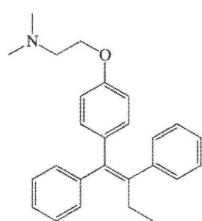
19 Fig. 4. Competitive binding curves of estradiol, tamoxifen, 4-hydroxytamoxifen,
20 RID-SB1 and RID-G. ER α competitive binding activity was measured as described in
21 Materials and methods. Estradiol, tamoxifen, 4-hydroxytamoxifen, RID-G and
22 RID-SB1 are represented as \circ , \diamond , \square , \blacksquare , \blacktriangle respectively.

23 Fig. 5. The scattergram of RIDs for the proliferation inhibition activity and ER α binding
24 activity. The *X*-axis (-MG-MID) represents the proliferation inhibition of forty-eight
25 RIDs in 39 cell lines. The *Y*-axis (-Log (IC₅₀)) represents the competitive ER α binding

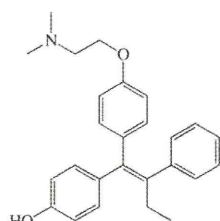
1 activity. Tamoxifen, RID-SB1 and SB8 are represented as \diamond , \square , \triangle , respectively.

Fig. 1

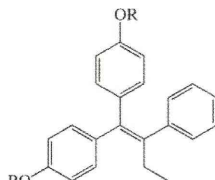
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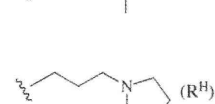
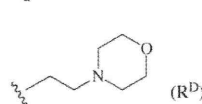
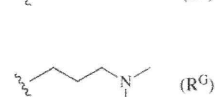
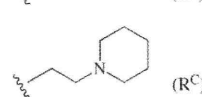
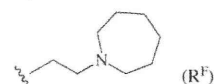
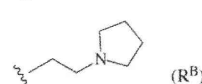
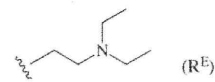
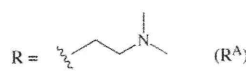
tamoxifen



4-hydroxytamoxifen

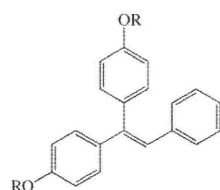


RID-X
(R = R^X)

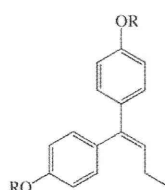


B

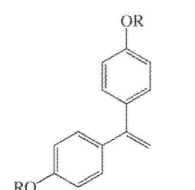
R = R^B
R^G



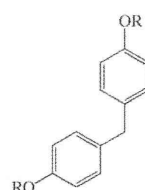
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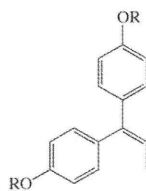
RID-SX2
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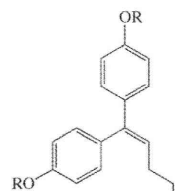
RID-SX3
(R = R^X)



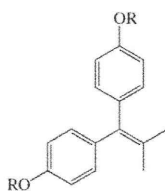
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(R = R^X)



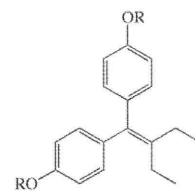
RID-SX7
(R = R^X)



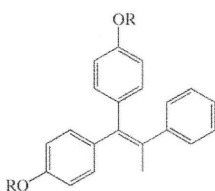
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(R = R^X)



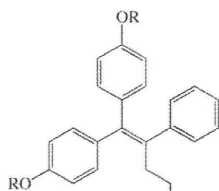
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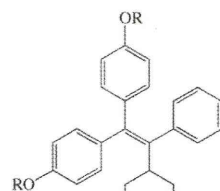
RID-SX10
(R = R^X)



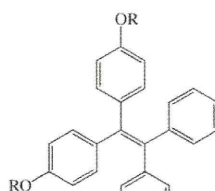
RID-SX11
(R = R^X)



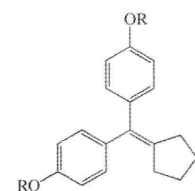
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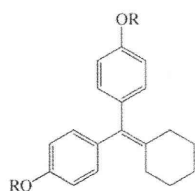
RID-SX13
(R = R^X)



RID-SX14
(R = R^X)



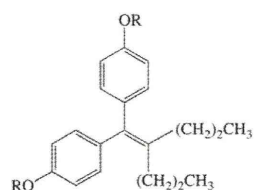
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(R = R^X)



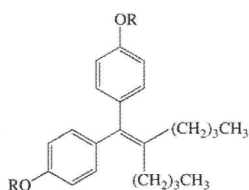
RID-SX16
(R = R^X)

C

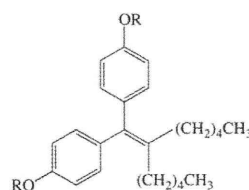
R = R^B
R^F
R^G



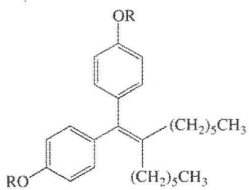
RID-SX22
(R = R^X)



RID-SX23
(R = R^X)



RID-SX17
(R = R^X)



RID-SX24
(R = R^X)

Fig. 2A

A

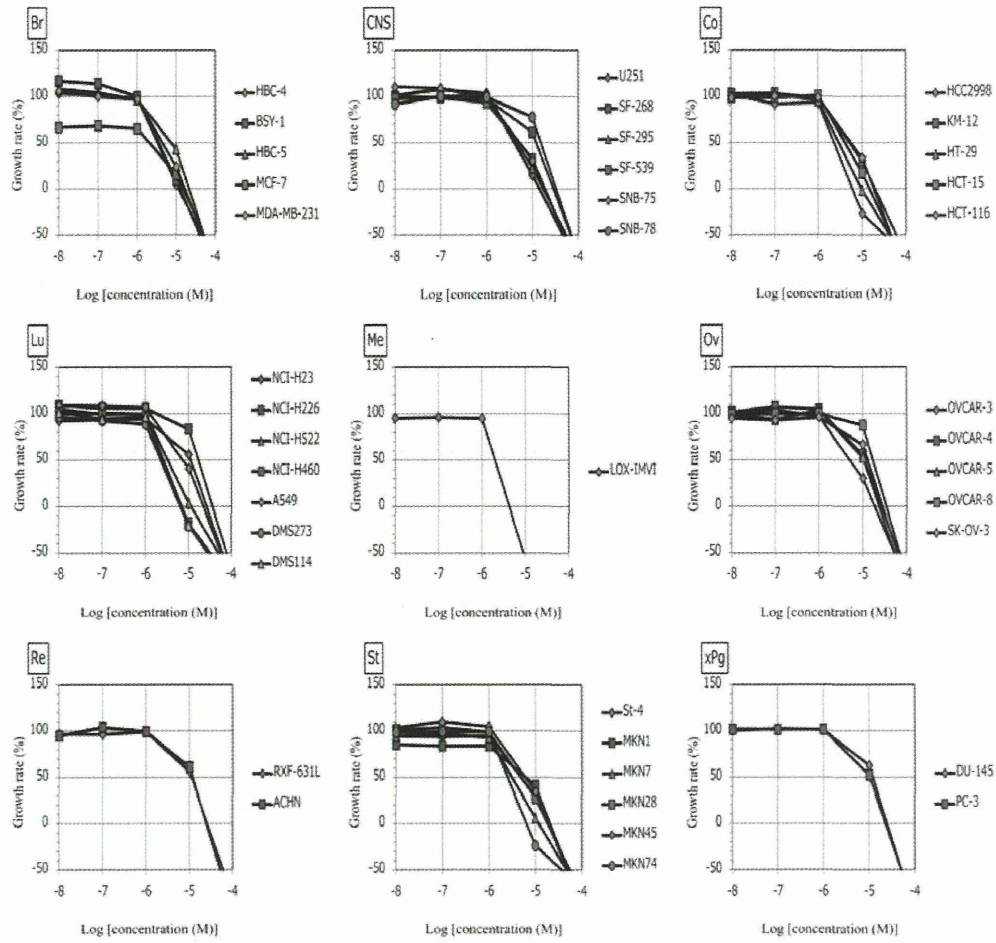


Fig. 2B

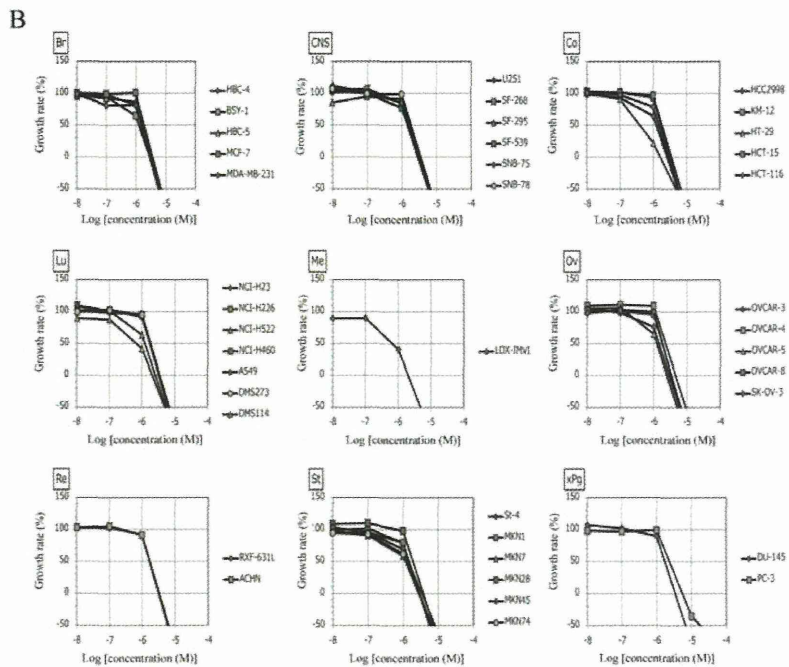


Fig. 3

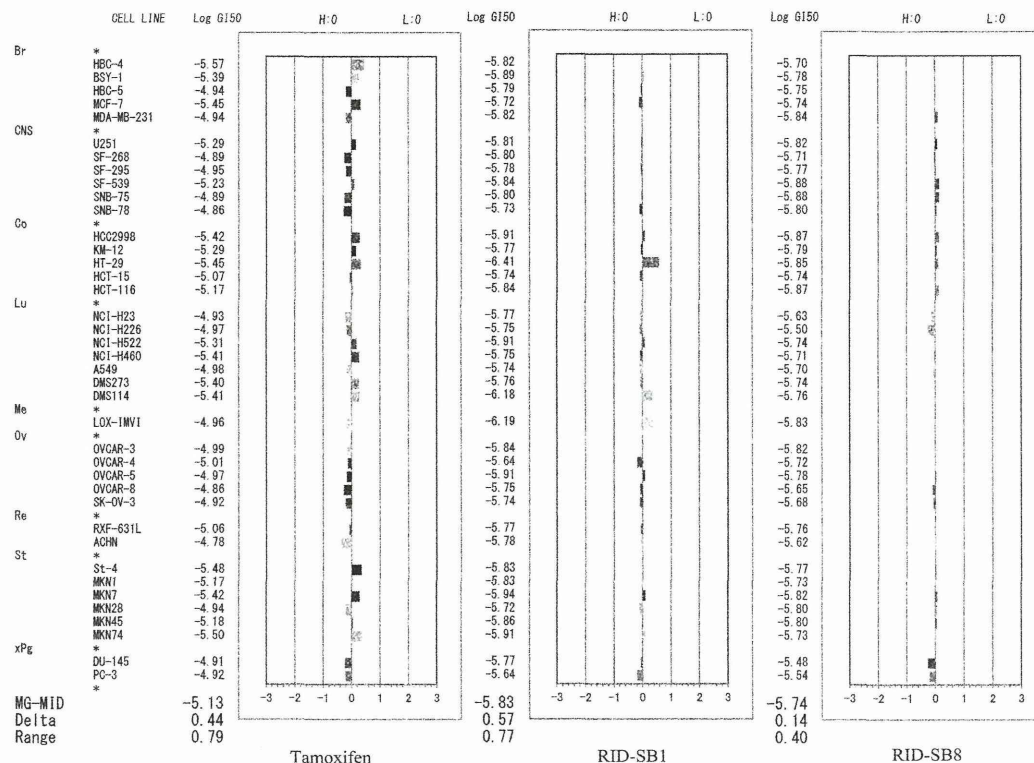


Fig. 4

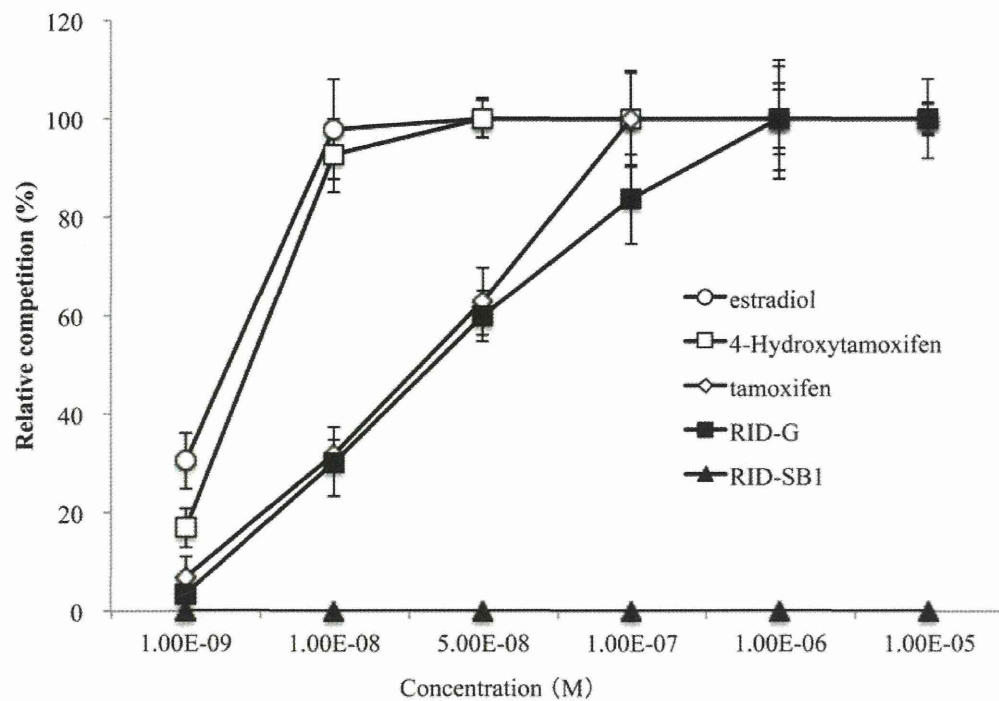


Fig. 5

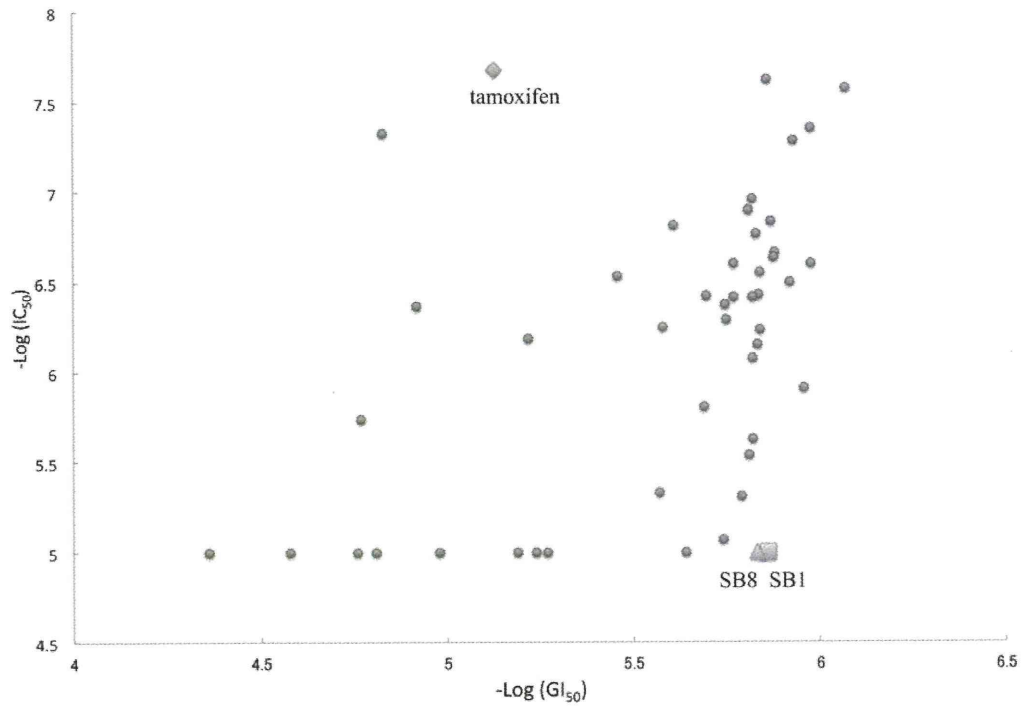


Table 1. Anti-tumor activity and ER α binding capacity of RIDs

Chemical	MG-MID (μ M)	ER α binding activity IC ₅₀ (nM)	Generation	Chemical	MG-MID (μ M)	ER α binding activity IC ₅₀ (nM)	Generation
17 β -estradiol	-	1.89		RID-SG2	15.50	>10000	2nd
4-Hydroxytamoxifen	5.50	2.72		RID-SG3	6.46	>10000	2nd
Tamoxifen	7.41	21.1		RID-SG4	26.3	>10000	2nd
RID-A	1.48	171	1st	RID-SG7	17.4	>10000	2nd
RID-B	1.17	52.4	1st	RID-SG8	1.55	2910	2nd
RID-C	3.47	295	1st	RID-SG9	1.82	8570	2nd
RID-D	14.8	47.7	1st	RID-SG10	1.45	372	2nd
RID-E	1.20	320	1st	RID-SG11	1.55	127	2nd
RID-F	2.45	154	1st	RID-SG12	1.38	23.9	2nd
RID-G	0.85	26.7	1st	RID-SG13	1.32	234	2nd
RID-H	1.05	44.1	1st	RID-SG14	1.51	109	2nd
RID-SB1	1.38	>10000	2nd	RID-SG15	1.51	835	2nd
RID-SB2	5.40	>10000	2nd	RID-SG16	1.10	1230	2nd
RID-SB3	10.5	>10000	2nd	RID-SB22	1.51	383	3rd
RID-SB4	43.7	>10000	2nd	RID-SB23	1.70	253	3rd
RID-SB7	5.77	>10000	2nd	RID-SB17	2.00	380	3rd
RID-SB8	1.82	>10000	2nd	RID-SB24	2.63	568	3rd
RID-SB9	2.69	4640	2nd	RID-SF22	1.79	421	3rd
RID-SB10	1.51	2370	2nd	RID-SF23	6.03	656	3rd
RID-SB11	1.44	582	2nd	RID-SF17	12.0	428	3rd
RID-SB12	1.34	145	2nd	RID-SF24	17.0	1840	3rd
RID-SB13	1.31	218	2nd	RID-SG22	1.05	253	3rd
RID-SB14	1.44	280	2nd	RID-SG23	1.70	387	3rd
RID-SB15	1.62	4880	2nd	RID-SG17	1.78	515	3rd
RID-SB16	1.46	702	2nd	RID-SG24	2.04	1560	3rd
RID-SG1	2.29	>10000	2nd				

Table 2. COMPARE Analysis of RID-SB1 and RID-SB8.

Chemical	Ranking	Drug or inhibitor	r	Function
RID-SB1	1	NVP-AEW541	0.505	IGF-1R inhibitor
	2	Bortezomib	0.481	proteasome inhibitor
	3	RDEA119	0.474	MEK inhibitor
	:	:	:	:
	>100	tamoxifen	0.199	ER antagonist
RID-SB8	1	PB28 dihydrochloride	0.563	sigma-2 receptor agonist
	2	MEK inhibitor I	0.549	MEK inhibitor
	3	Raf1 Kinase Inhibitor I	0.549	Raf1 inhibitor
	:	:	:	:
	>100	tamoxifen	0.314	ER antagonist

^a The ID for the chemical in JFCR39 database

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